

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/586,245
Source: IFWP
Date Processed by STIC: 7/26/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<<http://www.uspto.gov/efb/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)**
- 2. U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):**
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

10/586,245

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 Invalid Line Length The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.

- 4 ✓ Non-ASCII The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for **each** skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If **intentional**, please insert the following lines for **each** skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is **MANDATORY** if n's or Xaa's are present.
 In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.

- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence. (see item 11 below)

- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is **MANDATORY** if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules

- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 Misuse of n/Xaa "n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid



IFWP

RAW SEQUENCE LISTING

DATE: 07/26/2006

PATENT APPLICATION: US/10/586,245

TIME: 14:00:36

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\07262006\J586245.raw

*see item 4 on
Error Summary
sheet*

3 <110> APPLICANT: Consejo Superior de Investigaciones Cientificas
5 <120> TITLE OF INVENTION: GENERATION OF SPECIFIC ADHESION IN GRAM-NEGATIVE BACTERIA BY

MEANS

6 OF FIXING IMMUNOGLOBULIN SINGLE DOMAINS ON THEIR SURFACE WITH
7 AUTOTRANSPORTERS
9 <130> FILE REFERENCE: P1375PC
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/586,245
C--> 11 <141> CURRENT FILING DATE: 2006-07-11
11 <150> PRIOR APPLICATION NUMBER: ES P200400073
W--> 12 <151> PRIOR FILING DATE: 2004-01-14 ~~(January 14, 2004)~~
14 <160> NUMBER OF SEQ ID NOS: 10
16 <170> SOFTWARE: PatentIn version 3.1

*see
pg 1,3,5-6*

*delete 2004-01-14 is the
correct
format.*

ERRORED SEQUENCES

18 <210> SEQ ID NO: 1
19 <211> LENGTH: 5587
20 <212> TYPE: DNA
21 <213> ORGANISM: Artificial
W--> 23 <220> FEATURE:
23 <223> OTHER INFORMATION: DNA sequence of plasmid pVamyB
25 <400> SEQUENCE: 1

**Does Not Comply
Corrected Diskette Needed**
*insert
<220> whenever <221>, <222>, or <223>
is shown.*

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*<220>
is a header
only.
It never
has a
response*

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RAW SEQUENCE LISTING

DATE: 07/26/2006

PATENT APPLICATION: US/10/586,245

TIME: 14:00:36

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119 tcacatg 5587

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123 <211> LENGTH: 5563

124 <212> TYPE: DNA

125 <213> ORGANISM: Artificial

W--> 127 <220> FEATURE:

127 <223> OTHER INFORMATION: DNA sequence of plasmid pVLMB10a

OK-> 129 <400> SEQUENCE: 2

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176	tgaaaatcaa	tcaattcgag	attgtcccta	gtgcgggtat	ccgttacagc	cgctgtcat	2820
177	ctgcagatta	caagttgggt	gacgacagt	ttaaagtaag	ttctatggca	gtgaaaacac	2880
178	taacggccgg	actggatttt	gcttatcggt	ttaaagtcgg	caaccttacc	gtaaaacct	2940
179	tgttatctgc	agcttacttt	gccaattatg	gcaaagccgg	cgtgaatgtg	ggcggtaaat	3000
180	ccttcgccta	taaagcagat	aatcaacagc	aatattcagc	aggcgtcgcg	ttactgtacc	3060
181	gtaatgttac	attaaacgta	aatggcagta	ttacaaaagg	aaaacaattg	gaaaaacaaa	3120
182	aatccggaca	aattaaaata	cagattcggt	tctaaaatac	caaattcata	gcaaaaataa	3180
183	atgccgtctg	aactcaagct	tgacctgtga	agtgaaaaat	ggcgacacatt	gtgcgacatt	3240
184	ttttttgtct	gcggtttacc	gctactgcgt	cacggatccc	cacgcgccct	gtagcggcgc	3300
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186	agcgcgcgct	cctttcgctt	tcttcccttc	ctttctcgcc	acgttcgcgc	gctttccccg	3420
187	tcaagctcta	aatcggggca	tccctttagg	gttccgattt	agtgcctttac	ggcacctcga	3480
188	ccccaaaaaa	cttgattagg	gtgatgggtc	acgtagtggg	ccatcgccct	gatagacggt	3540
189	ttttcgccct	ttgacgttgg	agtccacggt	ctttaatagt	ggactcttgt	tccaaactgg	3600
190	aacaacactc	aaccttatct	cggctctatc	ttttgattta	taagggattt	tgccgatttc	3660
191	ggcctatttg	ttaaaaaatg	agctgattta	acaaaaat	aacgcgaatt	ttaacaaaat	3720
192	attaacgttt	acaatttcag	gtggcacttt	tccgggaaat	gtgcgcggaa	cccctatttg	3780
193	tttatttttc	taaatacatt	caaatatgta	tccgctcatg	tcgagacggt	gggtgaggtt	3840
194	ccaactttca	ccataatgaa	ataagatcac	taccgggcgt	attttttgag	ttatcgagat	3900
195	tttcaggagc	taaggaagct	aaaatggaga	aaaaaatcac	tggatatacc	accgttgata	3960
196	tatcccaatg	gcacgtgtaa	gaacattttg	aggcatttca	gtcagttgct	caatgtacct	4020

RAW SEQUENCE LISTING

DATE: 07/26/2006

PATENT APPLICATION: US/10/586,245

TIME: 14:00:36

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\07262006\J586245.raw

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197 ataaccagac cggttcagctg gatattacgg ccttttttaa gaccgtaaag aaaaataagc 4080
198 acaagtttta tccggccttt attcacattc ttgcccgcct gatgaatgct catccggagt 4140
199 tccgtatggc aatgaaagac ggtgagctgg tgatatggga tagtggtcac ccttgttaca 4200
200 ccgttttcca tgagcaaact gaaacgtttt catcgctctg gagtgaatac cagcagcatt 4260
201 tccggcagtt tctacacata tattcgcaag atgtggcgctg ttacggtgaa aacctggcct 4320
202 atttccctaa aggggtttatt gagaatatgt ttttcgtctc agccaatccc tgggtgagtt 4380
203 tcaccagttt tgatttaaac gtggccaata tggacaactt cttcgccccc gttttcacca 4440
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206 atgagtggca gggcgggggc taattttttt aaggcagtta ttggtgccct taaacgcctg 4620
207 gtgctacgcc tgaataagtg ataataagcg gatgaatggc agaaattcga aagcaaattc 4680
208 gacccggctg tccggttcagg gcagggtcgt taaatagccg cttatgtcta ttgctggtt 4740
209 accggtttat tgactaccgg aagcagtgtg accgtgtgct tctcaaatgc ctgaggccag 4800
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213 ggtttgtttg ccggatcaag agctaccaac tctttttccg aaggtaactg gcttcagcag 5040
214 agcgcagata ccaataactg tccttctagt gtagecgtag ttaggccacc acttcaagaa 5100
215 ctctgtagca ccgcctacat accctgctct gctaatcctg ttaccagtgg ctgctgccag 5160
216 tggcgataay ccggttctta ccgggtttga ctcaagacga tagttaccgg ataaggcgca 5220
217 gcggtcgggc tgaacggggg gttcgtgcac acagcccagc ttggagcgaa cgacctacac 5280
218 cgaactgaga tacctacagc gtgagctatg agaaagcgcc acgcttcccg aaggagagaa 5340
219 ggcggacagg tatccggtaa gcggcagggt cggaacagga gagcgcacga gggagcttcc 5400
220 agggggaaac gcctgggtatc tttatagtcc tgcggggttt cgccacctct gacttgagcg 5460
221 tcgatttttg tgatgctcgt cagggggggcg gagcctatgg aaaaacgcca gcaacgcggc 5520
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225 <210> SEQ ID NO: 3

226 <211> LENGTH: 47

227 <212> TYPE: DNA

228 <213> ORGANISM: Artificial

W--> 230 <220> FEATURE:

230 <223> OTHER INFORMATION: Primer VHHA1

232 <400> SEQUENCE: 3

233 ctatgcgggc cagccggcca tggtcaggt gcagctgggtg gagtctt

47

236 <210> SEQ ID NO: 4

237 <211> LENGTH: 21

238 <212> TYPE: DNA

239 <213> ORGANISM: Artificial

W--> 241 <220> FEATURE:

241 <223> OTHER INFORMATION: Primer GEN III-Rev

243 <400> SEQUENCE: 4

244 accctcatag ttagcgtaac g

21

247 <210> SEQ ID NO: 5

248 <211> LENGTH: 44

249 <212> TYPE: DNA

250 <213> ORGANISM: Artificial

W--> 252 <220> FEATURE:

252 <223> OTHER INFORMATION: Primer Linker-A48-VamyA

254 <400> SEQUENCE: 5

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/586,245

DATE: 07/26/2006

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Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\07262006\J586245.raw

255 ggcggtccga ctgctaactc tggacaggtg cagctggtgg agtc 44
 258 <210> SEQ ID NO: 6
 259 <211> LENGTH: 30
 260 <212> TYPE: DNA
 261 <213> ORGANISM: Artificial
 W--> 263 <220> **FEATURE:** *insert <220>*
 263 <223> OTHER INFORMATION: Primer Vamy-Not
 OK-> 265 <400> **SEQUENCE:** 6
 266 gagtcattct gcggccgctg aggagacggt 30
 269 <210> SEQ ID NO: 7
 270 <211> LENGTH: 60
 271 <212> TYPE: DNA
 272 <213> ORGANISM: Artificial
 W--> 274 <220> **FEATURE:** *insert <220>*
 274 <223> OTHER INFORMATION: Primer Linker-A48
 OK-> 276 <400> **SEQUENCE:** 7
 277 accccgtctc acaactccca ccaggttcca tccgcaggcg gtccgactgc taactctgga 60
 280 <210> SEQ ID NO: 8
 281 <211> LENGTH: 37
 282 <212> TYPE: DNA
 283 <213> ORGANISM: Artificial
 W--> 285 <220> **FEATURE:** *insert <220>*
 285 <223> OTHER INFORMATION: Primer Linker -A48-Vamy-eag1
 OK-> 287 <400> **SEQUENCE:** 8
 288 attactcgcc ggccggtacc ccgtctcaca actccca 37
 291 <210> SEQ ID NO: 9
 292 <211> LENGTH: 33
 293 <212> TYPE: DNA
 294 <213> ORGANISM: Artificial
 W--> 296 <220> **FEATURE:** *insert <220>*
 296 <223> OTHER INFORMATION: Primer VL1
 OK-> 298 <400> **SEQUENCE:** 9
 299 gagtcattct agaggagcct tttttttgga gat 33
 302 <210> SEQ ID NO: 10
 303 <211> LENGTH: 26
 304 <212> TYPE: DNA
 305 <213> ORGANISM: Artificial
 W--> 307 <220> **FEATURE:** *insert <220>*
 307 <223> OTHER INFORMATION: Primer VL2
 OK-> 309 <400> **SEQUENCE:** 10
 310 ctgagatgag tttttgttct gcggcc 26

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/586,245

DATE: 07/26/2006

TIME: 14:00:37

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\07262006\J586245.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
 L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
 L:12 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
 L:23 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
 L:25 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:1
 L:127 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
 L:129 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:2
 L:230 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
 L:232 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:3
 L:241 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
 L:243 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:4
 L:252 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
 L:254 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:5
 L:263 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
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 L:276 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:7
 L:285 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
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 L:307 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
 L:309 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:10